	w shington Kalog
	CRF _rrors Correct W, the STIC Syste is Branch CRF Processing Date: 18/200 Edited by: Verified by: Verified by: Verified by:
11 1	CRF Processing Date: 9770/200 Edited by: Verified by: (STIC s
	Changed the margins in cases where the sequence text was 'wrapped' down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was  the prior application data; or other
	Added the mandatory heading and subheadings for *Current Application Data*.
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the
	applicant placed a response below the subheading, this was moved to its appropriate place.
	applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:
-	
	Inserted colons after headings/subheadings. Headings edited included: , '
	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of files.
•	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:
	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of fil page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:
-	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of fil page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:
	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:
	Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of fil page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/830,810

DATE: 05/29/2001 TIME: 19:12:50

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\I830810.raw

```
3 <110> APPLICANT: Matzuk, Martin
             Pei, Wang
     9 <120> TITLE OF INVENTION: OVARY SPECIFIC GENES AND PROTEINS
    13 <130> FILE REFERENCE: P01925WOO / 09807797 / OTA 99-48
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/830,810
C--> 19 <141> CURRENT FILING DATE: 2001-04-27
    23 <150> PRIOR APPLICATION NUMBER: 60/106,020
    25 <151> PRIOR FILING DATE: 1998-10-28
    29 <160> NUMBER OF SEQ ID NOS: 15
    33 <170> SOFTWARE: PatentIn version 3.0
    37 <210> SEQ ID NO: 1
    39 <211> LENGTH: 1277
    41 <212> TYPE: DNA
    43 <213> ORGANISM: Mus musculus
    47 <400> SEQUENCE: 1
    48 aaggegggeg aggegeggga egeacecatg tteeeggega geaegtteea eeeetgeeeg
                                                                             60
    50 catcettate egeaggeeae caaageeggg gatggetgga ggtteggage caggggetge
                                                                            120
    52 cgacccgcgc cccctcctt cctccccggc tacagacagc tcatggccgc ggagtacqtc
                                                                            180
    54 gacagecace agegggeaca geteatggee etgetgtege ggatgggtee eeggteggte
                                                                            240
    300
    58 ctcgggcgcc gcacgctgca gcctgcaggg tgccgagcca gccccgacgc ccgatcgggt
                                                                            360
    60 tectgteaac eccgtggeea egeeggegee gggagateee egegateetg geagacegta
                                                                            420
    62 geocogttet egteegtgae ettetgtgge eteteeteet eaetggaggt tgegggagge
                                                                            480
    64 aggcaqacac ccacgaaggg agaggggagc ccggcatcct cggggacccg ggaaccggag
                                                                            540
    66 ccgagagagg tggccgcgag gaaagcggtc ccccagccgc gaagcgagga gggcgatgtt
                                                                            600
    68 caggetgeag ggeaggeegg gtgggageag cagecaceae eggaggaeeg gaacagtgtg
                                                                            660
    70 geggegatge agtetgagee tgggagegag gageeatgte etgeegeaga gatggeteag
                                                                            720
    72 gaccccggtg attcggatgc ccctcgagac caggcctccc cgcaaagcac ggagcaggac
                                                                            780
    74 aaggagcgcc tgcgtttcca gttcttagag cagaagtacg gctactatca ctgcaaggac
                                                                            840
    76 tgcaaaatcc ggtgggagag cgcctatgtg tggtgtgtgc agggcaccag taaggtgtta
                                                                            900
    78 cttcaaacag ttctgccgag tgtgtgagaa atcctacaac ccttacagag tggaggacat
                                                                            960
    80 cacctgtcaa agttgtaaaa gaactagatg tgcctgccca gtcagatttc gccacgtgga
                                                                           1020
    82 ccctaaacgc ccccatcggc aagacttgtg tgggagatgc aaggacaaac gcctgtcctg
                                                                           1080
    84 cgacagcacc ttcagcttca aatacatcat ttagtgagag tcgaaaacgt ttctgctaga
                                                                           1140
    86 tggggctaat ggaatggaca agtgagcttt ctcccctctt cacctcttcc ctttccaaat
                                                                           1200
    88 tottcatgac agacagtgtt acttggatat aaagcotgtg aataaaaggt attgcaaaca
                                                                           1260
                                                                           1277
    90 aaaaaaaaa aaaaaaa
    93 <210> SEQ ID NO: 2
    95 <211> LENGTH: 361
    97 <212> TYPE: PRT
    99 <213> ORGANISM: Mus musculus
    103 <400> SEOUENCE: 2
    105 Met Phe Pro Ala Ser Thr Phe His Pro Cys Pro His Pro Tyr Pro Gln
    106 1
                        5
                                            10
    108 Ala Thr Lys Ala Gly Asp Gly Trp Arg Phe Gly Ala Arg Gly Cys Arg
    111 Pro Ala Pro Pro Ser Phe Leu Pro Gly Tyr Arg Gln Leu Met Ala Ala
```

RAW SEQUENCE LISTING DATE: 05/29/2001 PATENT APPLICATION: US/09/830,810 TIME: 19:12:50

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\I830810.raw

112			35					40					45				
	Glu	Tyr		Asp	Ser	His	Gln		Ala	Gln	Leu	Met		Leu	Leu	Ser	
115		50					55	,				60					
117	Arg	Met	Gly	Pro	Arg	Ser	Val	Ser	Ser	Arg	Asp	Ala	Ala	Val	Gln	Val	
118	65		-			70				_	75					80	
120	Asn	Pro	Arg	Arg	Asp	Ala	Ser	Val	Gln	Cys	Ser	Leu	Gly	Arg	Arg	Thr	
121			_	_	85					90			_		95		
123	Leu	Gln	Pro	Ala	Gly	Cys	Arg	Ala	Ser	Pro	Asp	Ala	Arg	Ser	Gly	Ser	
124				100					105					110	_		
126	Cys	Gln	Pro	Arg	Gly	His	Ala	Gly	Ala	Gly	Arg	Ser	Pro	Arg	Ser	Trp	
127			115					120					125				
129	Gln	Thr	Val	Ala	Pro	Phe	Ser	Ser	Val	Thr	Phe	Cys	Gly	Leu	Ser	Ser	
130		130					135					140					
132	Ser	Leu	Glu	Val	Ala	Gly	Gly	Arg	Gln	Thr		Thr	Lys	Gly	Glu	Gly	
	145					150					155					160	
	Ser	Pro	Ala	Ser	Ser	Gly	Thr	Arg	Glu	Pro	Glu	Pro	Arg	Glu		Ala	
136					165					170					175		
	Ala	Arg	Lys		Val	Pro	Gln	Pro		Ser	Glu	Glu	Gly	Asp	Val	Gln	
139				180					185	_				190			
	Ala	Ala	-	Gln	Ala	Gly	Trp		Gln	Gln	Pro	Pro		Glu	Asp	Arg	
142			195					200				_	205		_	_	
	Asn		Val	Ala	Ala	Met		Ser	Glu	Pro	Gly		Glu	Glu	Pro	Cys	
145	_	210	- 1	<b>~</b> 1			215			<b>6</b> 1	70 .	220	<b>7</b>	<b>7</b> . 7		_	
		Ата	Ата	GIU	Met		GIN	Asp	Pro	стА		ser	Asp	ATa	Pro		
	225	Cla	71.7	C0.~	Dwo	230	Cor	Th x	C1	Cln	235	Tvc	C1,,	7~~	T 011	240	
151	ASP	GIII	Ala	ser	Pro 245	GIII	Ser	1111	GIU	250	Asp	пÃ2	GIU	Arg	255	Arg	
	Dha	Cln	Dha	Lou	Glu	Gln	Luc	Tur.	Cl v		Фулт	Hie	Cue	Luc		Cve	
154	1116	GIII	1116	260	Giu	OLII	цуз	ı yı	265	ı yı	- Y -	1113	Cys	270	лор	Cys	
	Lvs	Tle	Ara		Glu	Ser	Ala	Tur		Trn	Cvs	Val	Gln		Thr	Ser	
157	27.0		275	_	0.1.0	001		280			0,0		285	01,		001	
	Lvs	Val			Lys	Gln	Phe		Ara	Val	Cvs	Glu		Ser	Tvr	Asn	
160	-1-	290	- 1		1		295	-1-	5		- 4	300			- 2 -		
	Pro		Arg	Val	Glu	Asp	Ile	Thr	Cys	Gln	Ser	Cys	Lys	Arg	Thr	Arg	
	305	4	,			310			_		315	-	•			320	
165	Cys	Ala	Cys	Pro	Val	Arg	Phe	Arg	His	Val	Asp	Pro	Lys	Arg	Pro	His	
166	-		•		325			•		330	_		-	_	335		
168	Arg	Gln	Asp	Leu	Cys	Gly	Arg	Cys	Lys	Asp	Lys	Arg	Leu	Ser	Cys	Asp	
169	, -		_	340	_	_	_	_	345	_	_	_		350	_	_	
171	Ser	Thr	Phe	Ser	Phe	Lys	Tyr	Ile	Ile								,
172			355					360									
174	<210	)> SI	EQ II	ON C	: 3												
	5 <211> LENGTH: 1817																
	3 <212> TYPE: DNA																
	O <213> ORGANISM: Mus musculus																
	4 <400> SEQUENCE: 3																
	5 gtcacagett teceetgeee gaatatggtg atetgtetee attgteeaga teaggatgat 7 tetttagaag aagteacaga ggaatgetat teceeaceea eeeteeagaa eetggeaatt																
187	tctt	taga	aag a	aagto	cacaç	ga go	gaat	gctat	t to	ccca	ccca	ccct	cca	gaa d	cctg	gcaati	Ē.
189	caga	agtc	cac t	gag	ggat	ga go	gccti	ggc	c att	tct	gctc	tcad	egga	cct q	gccc	cagagt	<u>.</u>

60 120 180 RAW SEQUENCE LISTING DATE: 05/29/2001 PATENT APPLICATION: US/09/830,810 TIME: 19:12:50

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\I830810.raw

```
191 ctgttcccag taatttttga ggaggccttc actgatggat atatagggat cttgaaggcc
                                                                          240
193 atgatacctg tgtggccctt cccatacctt tctttaggaa agcagataaa taattgcaac
                                                                          300
195 ctggagactt tgaaggctat gcttgaggga ctagatatac tgcttgcaca aaaggttcaa
                                                                          360
197 accagtaggt gcaaactcag agtaattaat tggagagaag atgacttgaa gatatgggct
                                                                          420
199 ggatcccatg aaggtgaagg cttaccagat ttcaggacag agaagcagcc aattgagaac
                                                                          480
201 agtgctggct gtgaggtgaa gaaagaattg aaggtgacga ctgaagtcct tcgcatqaaq
                                                                          540
203 ggcagacttg atgaatctac cacatacttg ttgcagtggg cccagcagag aaaagattct
                                                                          600
205 attcatctat tetgtagaaa getactaatt gaaggettaa eeaaageete agtgatagaa
                                                                          660
207 atcttcaaaa ctgtacacgc agactgtata caggagctta tcctaagatg tatctgcata
                                                                          720
209 gaagagttgg cttttcttaa tccctacctg aaactgatga aaagtctttt cacactcaca
                                                                          780
211 ctagatcaca tcataggtac cttcagtttg ggtgattctg aaaagcttga tgaggagaca
                                                                          840
213 atatteaget tgatttetea aetteeeaca etceaetgte tecagaaact etatgtaaat
                                                                          900
215 gatgtccctt ttataaaagg caacctgaaa gaatacctca ggtgcctgaa aaagcccttg
                                                                          960
217 gagacacttt gcatcagtaa ctgtgacctc tcacagtcag acttggattg cctgccctat
                                                                         1020
219 tgcctgaata tttgtgaact caaacatctg catattagtg atatatattt atgtgattta
                                                                         1080
221 ctccttgagc ctcttggttt tctccttgag agagttggag ataccctgaa aaccctggaa
                                                                         1140
223 ttggattcat gttgtatagt ggactttcag ttcagtgcct tgctgcctgc cctaaqccaa
                                                                         1200
225 tgttctcacc tcagagaggt cactttctat gataatgatg tttctctgcc tttcttgaaa
                                                                         1260
227 acaacttcta caccacacag ccctgctgag tcagctgatc tatgagtgtt accctgccc
                                                                         1320
229 totagagtgo tatgatgaca gtggtgtaat actaacacac agattagaaa gtttttgtco
                                                                         1380
231 tgagcttctg gatatactga gagccaaaag acagctccat agtgtctcct ttcaaacaac
                                                                         1440
233 caaatgctct aaatgtggtg ggtgctacat ttatgatcgg catacccaat gttgccgttt
                                                                         1500
235 tgtggaacta ctataagctt gattgtgaaa ctgagaaata gaaacttagt attggggact
                                                                         1560
237 gatgaaatcc taagtgaatg tccactgcta aatggagcat gaaaatgtca atcacctaaa
                                                                         1620
239 agtotgagat acacaggaaa gtoaataact toototgago tggtgaatgg atgttgcato
                                                                         1680
241 tgtagaaagt atcaagcact tgtagtttga atgtgttaca atagaagcac cattttatga
                                                                         1740
243 gactggccca atctgttgac tgcatacaat aaatctgttg acttattaaa tttttaaaaa
                                                                         1800
245 aaaaaaaaa aaaaaaa
                                                                         1817
248 <210> SEQ ID NO: 4
250 <211> LENGTH: 426
252 <212> TYPE: PRT
254 <213> ORGANISM: Mus musculus
258 <400> SEQUENCE: 4
260 Met Val Ile Cys Leu His Cys Pro Asp Gln Asp Asp Ser Leu Glu Glu
261 1
                                        10
263 Val Thr Glu Glu Cys Tyr Ser Pro Pro Thr Leu Gln Asn Leu Ala Ile
                                     25
266 Gln Ser Leu Leu Arg Asp Glu Ala Leu Ala Ile Ser Ala Leu Thr Asp
269 Leu Pro Gln Ser Leu Phe Pro Val Ile Phe Glu Glu Ala Phe Thr Asp
                            55
                                                 60
272 Gly Tyr Ile Gly Ile Leu Lys Ala Met Ile Pro Val Trp Pro Phe Pro
                        70
                                            75
275 Tyr Leu Ser Leu Gly Lys Gln Ile Asn Asn Cys Asn Leu Glu Thr Leu
276
278 Lys Ala Met Leu Glu Gly Leu Asp Ile Leu Leu Ala Gln Lys Val Gln
279
                100
                                     105
281 Thr Ser Arg Cys Lys Leu Arg Val Ile Asn Trp Arg Glu Asp Asp Leu
282
            115
                                120
```

RAW SEQUENCE LISTING DATE: 05/29/2001
PATENT APPLICATION: US/09/830,810 TIME: 19:12:50

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\1830810.raw

```
284 Lys Ile Trp Ala Gly Ser His Glu Gly Glu Gly Leu Pro Asp Phe Arg
        130
                             135
                                                 140
287 Thr Glu Lys Gln Pro Ile Glu Asn Ser Ala Gly Cys Glu Val Lys Lys
288 145
                         150
                                             155
290 Glu Leu Lys Val Thr Thr Glu Val Leu Arg Met Lys Gly Arg Leu Asp
                    165
                                         170
293 Glu Ser Thr Thr Tyr Leu Leu Gln Trp Ala Gln Gln Arg Lys Asp Ser
                180
                                     185
296 Ile His Leu Phe Cys Arg Lys Leu Leu Ile Glu Gly Leu Thr Lys Ala
297
           · 195
                                 200
299 Ser Val Ile Glu Ile Phe Lys Thr Val His Ala Asp Cys Ile Gln Glu
                             215
302 Leu Ile Leu Arg Cys Ile Cys Ile Glu Glu Leu Ala Phe Leu Asn Pro
                         230
                                             235
305 Tyr Leu Lys Leu Met Lys Ser Leu Phe Thr Leu Thr Leu Asp His Ile
                    245
                                         250
308 Ile Gly Thr Phe Ser Leu Gly Asp Ser Glu Lys Leu Asp Glu Glu Thr
                260
                                     265
311 Ile Phe Ser Leu Ile Ser Gln Leu Pro Thr Leu His Cys Leu Gln Lys
            275
                                 280
314 Leu Tyr Val Asn Asp Val Pro Phe Ile Lys Gly Asn Leu Lys Glu Tyr
        290
                             295
                                                 300
317 Leu Arg Cys Leu Lys Lys Pro Leu Glu Thr Leu Cys Ile Ser Asn Cys
318 305
                        310
320 Asp Leu Ser Gln Ser Asp Leu Asp Cys Leu Pro Tyr Cys Leu Asn Ile
                    325
                                         330
323 Cys Glu Leu Lys His Leu His Ile Ser Asp Ile Tyr Leu Cys Asp Leu
                340
                                     345
                                                         350
326 Leu Leu Glu Pro Leu Gly Phe Leu Leu Glu Arg Val Gly Asp Thr Leu
            355
                                 360
                                                     365
329 Lys Thr Leu Glu Leu Asp Ser Cys Cys Ile Val Asp Phe Gln Phe Ser
        370
                            375
                                                 380
332 Ala Leu Leu Pro Ala Leu Ser Gln Cys Ser His Leu Arg Glu Val Thr
333 385
                        390
                                             3.95
335 Phe Tyr Asp Asn Asp Val Ser Leu Pro Phe Leu Lys Thr Thr Ser Thr
336
                    405
                                         410
338 Pro His Ser Pro Ala Glu Ser Ala Asp Leu
                420
341 <210> SEQ ID NO: 5
343 <211> LENGTH: 1018
345 <212> TYPE: DNA
347 <213> ORGANISM: Mus musculus
351 <400> SEQUENCE: 5
352 gccatattga ggacctgcag tagaggtgga acccatgact ggcagcgcaa acacagtgat
                                                                            60
354 aacagetgag etecaageaa ggaeeeagga eettgeetea eeacagaeat aatettteee
                                                                           120
356 cacaacacet ccaccaagec geeetgtaaa tegacatgag tegeeacage accageageg
                                                                           180
358 tgaccgaaac cacagcaaaa aacatgctct ggggtagtga actcaatcag gaaaagcaga
                                                                           240
360 cttgcacctt tagaggccaa ggcgagaaga aggacagctg taaactcttg ctcagcacga
                                                                           300
362 tetgeetggg ggagaaagee aaagaggagg tgaacegtgt ggaagteete teccaggaag
                                                                           360
```

RAW SEQUENCE LISTING DATE: 05/29/2001 PATENT APPLICATION: US/09/830,810 TIME: 19:12:50

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\I830810.raw

```
364 gcagaaaacc accaatcact attgctacgc tgaaggcatc agtcctgccc atggtcactg
                                                                        420
366 tgtcaggtat agagctttct cctccagtaa cttttcggct caggactggc tcaggacctg
                                                                        480
368 tgttcctcag tggcctggaa tgttatgaga cttcggacct gacctgggaa gatgacgagg
                                                                        540
370 aagaggagga agaggaggag gaagaggatg aagatgagga tgcagatata tcgctagagg
                                                                        600
372 agatacctgt caaacaagtc aaaagggtgg ctccccagaa gcagatgagc atagcaaaga
                                                                        660
374 aaaagaaggt ggaaaaagaa gaggatgaaa cagtagtgag gcccagccct caggacaaga
376 gtccctggaa gaaggagaaa tctacaccca gagcaaagaa gccagtgacc aagaaatgac
                                                                        780
378 ctcatcttag catcttctgc gtccaaggca ggatgtccag cagctgtgtt ttggtgcagg
                                                                        840
380 tqtccaqccc caccacccta qtctqaatqt aataaqqtqq tqtqqctqta accctqtaac
                                                                        900
382 ccagccctcc agtttccgga ggtttttggt gaagagcccc cagcaagttc gcctaqggcc
                                                                        960
1018
387 <210> SEO ID NO: 6
389 <211> LENGTH: 207
391 <212> TYPE: PRT
393 <213> ORGANISM: Mus musculus
397 <400> SEQUENCE: 6
399 Met Ser Arg His Ser Thr Ser Ser Val Thr Glu Thr Thr Ala Lys Asn
400 1
                   5
                                       10
402 Met Leu Trp Gly Ser Glu Leu Asn Gln Glu Lys Gln Thr Cys Thr Phe
405 Arg Gly Gln Gly Glu Lys Lys Asp Ser Cys Lys Leu Leu Ser Thr
406
408 Ile Cys Leu Gly Glu Lys Ala Lys Glu Glu Val Asn Arg Val Glu Val
409
411 Leu Ser Gln Glu Gly Arg Lys Pro Pro Ile Thr Ile Ala Thr Leu Lys
                       70
                                           75
414 Ala Ser Val Leu Pro Met Val Thr Val Ser Gly Ile Glu Leu Ser Pro
417 Pro Val Thr Phe Arg Leu Arg Thr Gly Ser Gly Pro Val Phe Leu Ser
               100
                                   105
420 Gly Leu Glu Cys Tyr Glu Thr Ser Asp Leu Thr Trp Glu Asp Asp Glu
                               120
           115
423 Glu Glu Glu Glu Glu Glu Glu Glu Asp Glu Asp Glu Asp Ala Asp
                           135
                                               140
426 Ile Ser Leu Glu Glu Ile Pro Val Lys Gln Val Lys Arg Val Ala Pro
                       150
                                           155
429 Gln Lys Gln Met Ser Ile Ala Lys Lys Lys Lys Val Glu Lys Glu Glu
                                       170
432 Asp Glu Thr Val Val Arg Pro Ser Pro Gln Asp Lys Ser Pro Trp Lys
433
               180
                                   185
435 Lys Glu Lys Ser Thr Pro Arg Ala Lys Lys Pro Val Thr Lys Lys
           195
                               200
438 <210> SEQ ID NO: 7
440 <211> LENGTH: 214
442 <212> TYPE: DNA
444 <213> ORGANISM: Mus musculus
448 <400> SEQUENCE: 7
449 acagcagagg tgatgctcag aaatcaagtt ttaacagagg gccaggtgct tctagagtag
                                                                        60
                                                                        120
451 gaggggattg cacacetece cacecetee tettteecag gettettaae ageetgetgt
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/830,810

DATE: 05/29/2001 TIME: 19:12:51

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05292001\I830810.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application Number

L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date

PCT

RAW SEQUENCE LISTING DATE: 05/18/2001 PATENT APPLICATION: US/09/830,810 TIME: 11:00:50

Input Set : A:\P01925US1 seq list.txt
Output Set: N:\CRF3\05182001\I830810.raw

	4		APPLICANT: Matzuk, Martin Pei, Wang TITLE OF INVENTION: OVARY SPECIFIC GENES AND PROTEINS	Does Not Comply
			FILE REFERENCE: P01925W00 / 09807797 / OTA 99-48	Does Not Company Corrected Diskette Needed
C>	10	<140>	CURRENT APPLICATION NUMBER: US/09/830,810	001103
C>	11	<141>	CURRENT FILING DATE: 2001-04-27	
	13	<150>	PRIOR APPLICATION NUMBER: 60/106,020	
	14	<151>	PRIOR FILING DATE: 1998-10-28	
	16	<160>	NUMBER OF SEC ID NOS: 15	

## ERRORED SEQUENCES

18 <170> SOFTWARE: PatentIn version 3.0

	495	<210> SEQ ID NO: 15	
		<211> LENGTH: 364	
	497	<212> TYPE: DNA	
	498	<213> ORGANISM: Mus musculus	
	500	<400> SEQUENCE: 15	
	501	ctcttatctg cacaggagaa atctacaccc agagcaaaga agccagtgac caagaaatga	60
	503	cctcatctta gcatcttctg cgtccaaggc aggatgtcca gcagctgtgt tctggtgcag	120
	505	gtgtccagcc ccaccaccct agtctgaatg taataaggtg gtgtggctgt aaccctgtaa	180
	507	cccagccctc cagtttccgg aggtttttgg tgaagagccc ccagcaagtt cgcctagggc	240
	509	cacaataaaa tttgcatgat caggacctcc ctctgcctcc ccctccctgg atgggtctcc	300
	511	togotgotgo gatagotoat gtgoccagoa gagggoaaco acgagoaaga aaccagocco	360
	513	atgt	364
E>	523/	25022143.1 Page 1	

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/830,810

DATE: 05/18/2001 TIME: 11:00:51

Input Set : A:\P01925US1 seq list.txt
Output Set: N:\CRF3\05182001\I830810.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:523 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:369 SEQ:15 L:523 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:12 L:523 M:252 E: No. of Seq. differs, <211>LENGTH:Input:364 Found:369 SEQ:15